

PATENT APPLICATION: US/09/901,484A

DATE: 03/05/2002 TIME: 10:08:34

Input Set : D:\GEN-T111XC3D2.ST25.txt
Output Set: N:\CRF3\03052002\I901484A.raw

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3 -: 110> APPLICANT: Cohen, Daniel
         Blumenfeld, Marta
 4
         Chumakov, Ilya
         Bougueleret, Lydie
8 -120> TITLE OF INVENTION: Prostate Cancer Gene
10 -(130) FILE REFERENCE: GEN-T111XC3D2
12 -: 140 > CURRENT APPLICATION NUMBER: US 09/901,484A
13 -: 141> CURRENT FILING DATE: 2001-07-09
15 -(150) PRIOR APPLICATION NUMBER: US 08/996,306
16 <1151> PRIOR FILING DATE: 1997-12-22
18 -: 150> PRIOR APPLICATION NUMBER: US 60/099,658
19 -: 151> PRIOR FILING DATE: 1998-09-09
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22 -: 151: PRIOR FILING DATE: 1998-12-22
24 -(150) PRIOR APPLICATION NUMBER: US 09/338,907
25 +1151> PRIOR FILING DATE: 1999-06-23
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28 -(151> PRIOR FILING DATE: 2001-05-11
30 <160> NUMBER OF SEQ ID NOS: 578
32 (170) SOFTWARE: PatentIn version 3.1
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35 -: 211> LENGTH: 56516
36 + 12120 + TYPE: DNA
37 -1.13> ORGANISM: Homo sapiens
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40 DEED NAME/KEY: promoter
41 HPD - LCCATION: (1629)..(1870)
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(0.D.D LOCATION: (1998)..(2000)
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52 0001 → NAME/KEY: misc_feature
53 < ADD LOCATION: (2001)..(2216)
54 <:23> OTHER INFORMATION: exon 1
57 KUDD> FEATURE:
רם ייספיי, אָאַאַטיילעטיי
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ENTERED

D. SAMBYERY of Local test the
Public Location: (11694)...(14337)

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- 66 <223> OTHER INFORMATION: Tyr Phos
- 69 (220> FEATURE:
- 70 (221) NAME/KEY: primer\_bind
- 71 + 2225 LOCATION: (11930)..(11947)
- 72 (223) OTHER INFORMATION: upstream amplification primer 4-77 SEQ ID 42
- 75 (220) FEATURE:
- 76 -: 221 NAME/KEY: allele
- 77 -: 222: LOCATION: (12057)..(12103)
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- 81 2200- FEATURE:
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- 83 222 LOCATION: (12339)..(12358)
- 84 223 OTHER INFORMATION: downstream amplification primer 4-77 SEQ ID 51, complement
- 37 2200 FEATURE:
- 88 1221 NAME/KEY: primer\_bind
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- 90 +223 OTHER INFORMATION: upstream amplification primer 4-73 SEQ ID 64
- 93 -(220 FEATURE:
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- 96 1223 OTHER INFORMATION: polymorphic fragment 4-73 SEQ ID 58
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- 102 (223) OTHER INFORMATION: downstream amplification primer 4-73 SEQ ID 67, complement
- 105 220> FEATURE:
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- 107 (222) LOCATION: (17997)..(18003)
- 108  $\pm 223$  OTHER INFORMATION: n = a, c, g, or t.
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- 112 +0.221 NAME/KEY: misc\_feature
- 113 -(222) LOCATION: (18196)..(18265)
- 114  $\pm (223)$  OTHER INFORMATION: exon 2
- 117 1220 > FEATURE:
- 118 MAME/KEY: misc\_feature
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- 131 < 222> LOCATION: (25571)..(25660)
- 132 (223> OTHER INFORMATION: exon 4
- 135 <230> FEATURE:

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108 <111> NAME/KEY: allele

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Input Set : D:\GEN-T111XC3D2.ST25.txt
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			catatttact				240							
			togcacactg				300							
			agttgtgcac				360							
			tacagtacag				420							
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			aaggtcatca				540							
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			aggtttccat				660							
			tatatatatg				720							
			catattaaac				780							
			tctgctttaa				840							
			aagtatgctt				900							
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			caagagtgaa				1020							
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268	geccaggtge	ctgcaagaat	tagacctccg	ataacgttaa	cacccacttt	ctcactgctc	1320							
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			accgcgcgca				1440 1500							
			agcacccacc				1500							
			ggcccagctg				1500							
			ctgtactgca				1630							
			ggaactatgg				1740							
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			toogggaggg				1860							
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312	eggagaegte	tacactccga	agtgcgcttt	tacagtgacc	tcttgaaaca	gaagtacaat	2640
314	teggtettgt	gttettteee	ctggacaagt	gaaagctggg	cgaagaaatg	aatacatttg	2700
316	ttaaccgtag	aageetaaet	agatacaatt	cttgccaact	ttaactgggc	ttgaatgtgt	2760
318	gggtgatctg	ttgtctgatt	actttctttc	tgttactgtt	tototgtaga	gattggattc	2820
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324	getttattet	tggttattcc	taatgtccac	ctagtcctct	tttacttttc	ttggtagggt	3000
		tggggaaatg					3060
		ctgtttttac					3120
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		ggctaatttt					3360
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		gcaagttctt					3600
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		catagaattt					4.54
		tggcaggtaa					463
		astittisti					468 131
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		gaatototoa					
		ataagtaata					4860 1000
		tocagttaag					4939 4980
		tgtgctgcca					4980 5040
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3 11		ggettgggge					5.5 5110
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## VERIFICATION SUMMARY

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Input Set : D:\GEN-T111XC3D2.ST25.txt
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